

09/973, 994

4/04/05

GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: June 15, 2003, 02:47:25 ; Search time 971.806 Seconds
(without alignments)
9193.767 Million cell

updates/sec

Title: US-09-973-994-79
Perfect score: 307
Sequence: 1 ggtgcgatcctagaattgca.....gccatcactcggatcgcacc 307

Scoring table: OLIGO_NUC
Gapop 60.0 , Gapext 60.0

Searched: 2054640 seqs, 14551402878 residues

Word size : 0

Total number of hits satisfying chosen parameters: 4109280

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

Database : GenEmbl:*
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2: gb_htg:*
3: gb_in:*
4: gb_om:*
5: gb_ov:*
6: gb_pat:*
7: gb_ph:*
8: gb_pl:*
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11: gb_sts:*
12: gb_sy:*
13: gb_un:*
14: gb_vi:*
15: em_ba:*
16: em_fun:*
17: em_hum:*
18: em_in:*
19: em_mu:*
20: em_om:*
21: em_or:*
22: em_ov:*
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26: em_ro:*
27: em_sts:*
28: em_un:*
29: em_vi:*

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30: em_htg_hum:*
31: em_htg_inv:*
32: em_htg_other:*
33: em_htg_mus:*
34: em_htg_pln:*
35: em_htg_rod:*
36: em_htg_mam:*
37: em_htg_vrt:*
38: em_sy:*
39: em_htgo_hum:*
40: em_htgo_mus:*
41: em_htgo_other:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Match	Query Length	DB	ID	Description

1	22	7.2	131427	2	AF121897	AF121897 Homo
sapi						
2	22	7.2	185982	2	AC073231	AC073231 Homo
sapi						
3	22	7.2	340000	9	HS21C080	AL163280 Homo
sapi						
4	21	6.8	185775	2	AL772276	AL772276 Mus
muscu						
5	20	6.5	65711	2	AC117583	AC117583 Mus
muscu						
6	20	6.5	87971	2	AC117094	AC117094
Rattus no						
7	20	6.5	94820	2	AC119140	AC119140
Rattus no						
c 8	20	6.5	140466	2	AC095152	AC095152
Rattus no						
9	20	6.5	141017	2	AC116962	AC116962
Dictyoste						
c 10	20	6.5	151374	2	AC111772	AC111772
Rattus no						
11	20	6.5	274947	2	AL731766	AL731766 Mus
muscu						
12	19	6.2	1627	8	ZMKN1	X61308 Z.mays
Knot						
13	19	6.2	5200	1	AY005468	AY005468
Pseudoalt						
c 14	19	6.2	33184	3	U70856	U70856
Caenorhabdi						
15	19	6.2	70162	2	AC026568	AC026568 Homo
sapi						
c 16	19	6.2	85458	2	AC106054	AC106054
Rattus no						
17	19	6.2	89207	2	AC124098	AC124098 Mus
muscu						
c 18	19	6.2	104600	9	AP001166	AP001166 Homo

sapi						
c 19	19	6.2	108400	9	AC005193	AC005193 Homo
sapi						
c 20	19	6.2	110000	2	AC129176_2	Continuation
(3 of						
21	19	6.2	123277	9	AC005016	AC005016 Homo
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c 23	19	6.2	134979	9	AL359963	AL359963
Human DNA						
24	19	6.2	135607	2	AC115922	AC115922 Mus
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c 25	19	6.2	136283	9	AP005202	AP005202 Homo
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c 27	19	6.2	140702	2	AC068420	AC068420 Homo
sapi						
28	19	6.2	141866	9	AL451062	AL451062
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29	19	6.2	143595	2	AC027563	AC027563 Homo
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30	19	6.2	148083	9	AF111170	AF111170 Homo
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32	19	6.2	155711	3	VYIVD10	AL360354
Plasmodiu						
c 33	19	6.2	156471	2	AC013568	AC013568 Homo
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34	19	6.2	162743	4	AC091728	AC091728 Bos
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c 35	19	6.2	162810	9	AC006382	AC006382 Homo
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36	19	6.2	162900	2	AC123889	AC123889
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Rattus no						
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sapi						
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muscu						
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muscu						
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muscu						

LOCUS PIAEMB2R 861 bp mRNA linear PLN 16-JUN-1999
 DEFINITION *Picea glauca* late embryogenesis abundant protein (EMB2) mRNA, complete cds.
 ACCESSION L47115
 VERSION L47115.1 GI:1350521
 KEYWORDS late embryogenesis abundant protein.
 SOURCE *Picea glauca* (white spruce)
 ORGANISM *Picea glauca*
 Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Coniferopsida; Coniferales; Pinaceae; *Picea*.
 REFERENCE 1 (bases 1 to 861)
 AUTHORS Dong, J.Z. and Dunstan, D.I.
 TITLE Cloning and characterization of six embryogenesis-associated cDNAs from somatic embryos of *Picea glauca* and their comparative expression during zygotic embryogenesis
 JOURNAL Plant Mol. Biol. 39 (4), 859-864 (1999)
 PUBMED 10350098
 REFERENCE 2 (bases 1 to 861)
 AUTHORS Dong, J.Z. and Dunstan, D.I.
 TITLE Direct Submission
 JOURNAL Submitted (30-MAY-1996) Crop Science, Horticulture Research International, East Malling, West Malling, Kent ME19 6BJ, UK
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 /note="(vector lambda zap)"
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 SKEVDEINAKFAEAREELEMEMESKETVYFDEEAETARDAVKQTLEMFDALLTKLPEA
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 121 catccaaccc ttaacctccg attctcaagc ccgctgaaaa accccatcaa gtcaacccta
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841 taagcttatc ataatttctt t

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LOCUS PIAEMB6R 942 bp DNA linear PLN 16-JUN-1999

DEFINITION *Picea glauca* late embryogenesis abundant protein (EMB6) mRNA, complete cds.

ACCESSION L47116

VERSION L47116.1 GI:1350540

KEYWORDS late embryogenesis abundant protein.

SOURCE *Picea glauca* (white spruce)

ORGANISM *Picea glauca*
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Coniferopsida; Coniferales; Pinaceae; *Picea*.

REFERENCE 1 (bases 1 to 942)

AUTHORS Dong, J.Z. and Dunstan, D.I.

TITLE Cloning and characterization of six embryogenesis-associated cDNAs from somatic embryos of *Picea glauca* and their comparative expression during zygotic embryogenesis

JOURNAL Plant Mol. Biol. 39 (4), 859-864 (1999)

PUBMED 10350098

REFERENCE 2 (bases 1 to 942)

AUTHORS Dong, J.Z. and Dunstan, D.I.

TITLE Direct Submission

JOURNAL Submitted (30-MAY-1996) Crop Science, Horticulture Research International, East Malling, West Malling, Kent ME19 6BJ, UK

FEATURES

Location/Qualifiers

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/db_xref="taxon:3330"
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5'UTR <1..49
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CDS 50..769
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polyA_site 942
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421 taatgaaacg catttttggt ttaaaacccc ggatcctgtg cggagacctg atcgttcgct
481 atttttaaat caaagccatg caaattccgc ttccggaagt gatagtccat ctcgatggcc
541 tgcaacgccg aaggcaccga aacgtgtagt gaagcatcgg aatcttctcg cgagtataa

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661 tgttgttgat gcagagcata ctatttgctt gaagccgggc attgctacaa acagggcttt
721 atttagcacg gatgagcttt ctggggcatt gtcccgtcac tgcatataga aagtttacga
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901 taaactctgt aaaaacaaat cttcaaccag ttttggggtc cc

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